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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Fri Jun 08 19:25:03 EDT 2007

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Reviewer Comments:

<210> 13
<211> 21
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<213> Artificial Sequence
<220>
<221> primer_bind
<223> reverse primer specific for TCR BV3 used in real-time PCR analysis
<400> 13
ggtgctggcg gactccagaa t 21

The above <213> Artificial Sequence is in an incorrect position; all numeric identifiers must be directly under each other. Do not use Tab keys. Same type of error in Sequences 20, 43, 50, 53, 68.

<400> 168
tacttctgtg ccagcagttc cctcgctact gctgaagctt tccttggaca aggc 54
??

??

??

??

Please delete the ?'s at the end of the submitted file.

Application No: 10612468 Version No: 2.0

Input Set:

Output Set:

Started: 2007-06-07 09:18:35.574
Finished: 2007-06-07 09:18:39.094
Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 520 ms
Total Warnings: 116
Total Errors: 118
No. of SeqIDs Defined: 168
Actual SeqID Count: 168

Error code	Error Description
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E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (1)
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Input Set:

Output Set:

Started: 2007-06-07 09:18:35.574
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Total Errors: 118
No. of SeqIDs Defined: 168
Actual SeqID Count: 168

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Input Set:

Output Set:

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Finished: 2007-06-07 09:18:39.094
Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 520 ms
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Total Errors: 118
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Actual SeqID Count: 168

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E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (25) This error has occurred more than 20 times, will not be displayed
E 249	Order Sequence Error <211> -> <213>; Expected Mandatory Tag: <212> in SEQID (146)
E 250	Structural Validation Error; Sequence listing may not be indexable

SEQUENCE LISTING

<110> Zhang, Jingwu Z.
Ho, Walter Kowk Keung
Zhang, Dongqing
Sun, Wei

<120> T Cell Receptor CDR3 Sequence and Methods for
Detecting and Treating Rheumatoid Arthritis

<130> D6622

<140> US 10/612,468
<141> 2003-07-02

<160> 168

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<223> part of the complementary determining region-3 (CDR3)
in the V(16 family (BV16 gene) of T cell receptors
(TCR) in patients with rheumatoid arthritis (RA)

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agtccgggg gcagtcgtt c 21

<210> 3
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<220>
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<223> conserved amino acid sequence derived from CDR3 of
TCR beta-chain BV16 in patients with RA

<400> 3
Ser Gln Ala Asp Gly Thr His

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Ser Ser Gly Gly Ser Leu Phe
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Ser Trp Gly Gly

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Ala Gly Pro Leu Glu Ala Gln Val Thr Gln Asn Pro Arg Tyr Leu
20 25 30
Ile Thr Val Thr Gly Lys Lys Leu Thr Val Thr Cys Ser Gln Asn
35 40 45
Met Asn His Glu Tyr Met Ser Trp Tyr Arg Gln Asp Pro Gly Leu
50 55 60
Gly Leu Arg Gln Ile Tyr Tyr Ser Met Asn Val Glu Val Thr Asp
65 70 75
Lys Gly Asp Val Pro Glu Gly Tyr Lys Val Ser Arg Lys Glu Lys
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Ser Leu Tyr Phe Cys Ala Ser Ser
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<210> 7
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<213> Homo sapiens

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Asp	Asn	Leu	Tyr	Trp	Tyr	Arg	Arg	Val	Met	Gly	Lys	Glu	Ile	Lys
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PCR analysis

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<210> 11
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